

W098/18930 SEQ ID 66

Qury Match 80.9%; Score 575.5; DB 4; Length 763;
Best Local Similarity 80.6%; Pred. No. 4,3e-56;
Matches 103; Conservative 18; Missmatches 9; Indels 1; Gaps 1;
Query Match 80.9%; Score 575.5; DB 4; Length 763;
Best Local Similarity 79.6%; Pred. No. 2,7e-56;
Matches 103; Conservative 18; Missmatches 9; Indels 1; Gaps 1;
Qy 7 CAYALNQHRS-GENKDNRNSVYVGSSOSSENTPDVSQSKERGIAEVIKTDGGY 65
Db 1 CSYELGXRHOGVKRSNSYDGGAGKAEINTPDESKERGIAEVIKTDGGY 60
Qy 66 VTSHGDDHYHYNKGKVYDALLPSEELLMKDNPYQLKDSDIVNEKGKVYVLLK 125
Db 61 VTSHGDDHYHYNKGKVYDALLPSEELLMKDNPYQLKDSDIVNEKGKVYVLLK 120
Qy 126 DAHAADNVRTK 136
Db 121 DAHAADNVRTK 131

W098/18930 SEQ ID 56

Qury Match 80.7%; Score 574; DB 4; Length 796;
Best Local Similarity 80.6%; Pred. No. 4,3e-56;
Matches 104; Conservative 16; Missmatches 9; Indels 0; Gaps 0;
Qy 8 AVYALNQHRSGENKDNRNSVYVGSSOSSENTPDVSQSKERGIAEVIKTDGGY 67
Db 1 SYELGXRHOGARTVKENRNSVYDGGAGTQTRNLTPESKERGIAEVIKTDGGY 60
Qy 68 SHGDHYHYNKGKVYDALLPSEELLMKDNPYQLKDSDIVNEKGKVYVLLDA 127
Db 61 SHGDHYHYNKGKVYDALLPSEELLMKDNPYQLKDSDIVNEKGKVYVLLDA 120
Qy 128 AHADNVRTK 136
Db 121 AHADNVRTK 129

1098/18930 SEQ ID 66

Qy 8 AVALNOHRSQENKNDRNNSVYDGSQSSKSENLTPEDESKREGINABQVIKITDGGVFT 67
Db 1 SYELGLYQARTVKNRNSVYDGSQATQITENLTPEDESKREGINABQVIKITDGGVFT 60
Qy 68 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Db 61 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Qy 128 AHDNVRTRK 136
Db 121 AHDNVRTRK 129

Matches 104; Conservative 16; Missmatches 9; Indels 0; Gaps 0;
Best Local Similarity 80.7%; Score 574; DB 4; Length 796;

Querty Match

Qy 7 CAYALNOHRS-GENKDNRNSVYDGSQSSKSENLTPEDESKREGINABQVIKITDGGVFT 65
Db 1 CSYELGRHOGVKESNSRVSYDGSQATQITENLTPEDESKREGINABQVIKITDGGVFT 60
Qy 66 VTSCHGDDHYHYNNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 125
Db 61 VTSCHGDDHYHYNNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 120
Qy 126 DAHADNVRTRK 136
Db 121 DAHADNVRTRK 131

Matches 103; Conservative 18; Missmatches 9; Indels 1; Gaps 1;
Best Local Similarity 79.6%; Pred. No. 2.7e-56;

Querty Match

1098/18930 SEQ ID 56

Qy 8 AVALNOHRSQENKNDRNNSVYDGSQSSKSENLTPEDESKREGINABQVIKITDGGVFT 67
Db 1 SYELGLYQARTVKNRNSVYDGSQATQITENLTPEDESKREGINABQVIKITDGGVFT 60
Qy 68 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Db 61 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Qy 128 AHDNVRTRK 136
Db 121 AHDNVRTRK 129

Matches 104; Conservative 16; Missmatches 9; Indels 0; Gaps 0;
Best Local Similarity 80.6%; Score 574; DB 4; Length 796;

Querty Match

Qy 8 AVALNOHRSQENKNDRNNSVYDGSQSSKSENLTPEDESKREGINABQVIKITDGGVFT 67
Db 1 SYELGLYQARTVKNRNSVYDGSQATQITENLTPEDESKREGINABQVIKITDGGVFT 60
Qy 68 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Db 61 SGCHHYXNGKPVYDALSSELMKPDNVQLKDAVINENVKGYIIKVGDGGVFT 60
Qy 128 AHDNVRTRK 136
Db 121 AHDNVRTRK 129

Matches 104; Conservative 16; Missmatches 9; Indels 0; Gaps 0;
Best Local Similarity 80.7%; Score 574; DB 4; Length 796;

Querty Match

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